

GenCore version 6.3

Copyright (c) 1993 - 2009 Bioceleration Ltd.

OM nucleic - nucleic search, using sw model

Run on: June 26, 2009, 03:19:14 ; Search time 752 Seconds
(without alignments)
14891.617 Million cell

updates/sec

Title: US-09-300-482-298
Perfect score: 388
Sequence: 1
ggagaaagaaagaaaagatg.....ttaagcctgggacccccgt 388

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 41078765 seqs, 14427166270 residues

Total number of hits satisfying chosen parameters: 82157530

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Published_Applications_NA_Main:*
1:
/ABSS/Data/CRF/ptodata/1/pubpna/US07_PUBCOMB.seq:*
2:
/ABSS/Data/CRF/ptodata/1/pubpna/US08_PUBCOMB.seq:*
3:
/ABSS/Data/CRF/ptodata/1/pubpna/US09A_PUBCOMB.seq:*
4:
/ABSS/Data/CRF/ptodata/1/pubpna/US09B_PUBCOMB.seq:*
5:
/ABSS/Data/CRF/ptodata/1/pubpna/US09C_PUBCOMB.seq:*
6:
/ABSS/Data/CRF/ptodata/1/pubpna/US09D_PUBCOMB.seq:*
7:
/ABSS/Data/CRF/ptodata/1/pubpna/US10A_PUBCOMB.seq:*
^